

Figure 1

```
1  gatcgcgaggagtcggtgcttttagtacgccgctggcacctttactctcgccggccgcgcgaa  60
                                     M S L
61  cccgtttgagctcggtatcctagtgcacacgcctttgcaagcgacggcgccatgagtcctg  120
    T S S S S V R V E W I A A V T I A A G T
121 acttccagttccagcgctacgagttgaatggatcgcagcagttaccattgctgctgggaca  180
    A A I G Y L A Y K R F Y V K D H R N K A
181 gctgcaattgggttatctagcttacaaaagattttatgttaaagatcatcgaaataaagct  240
    M I N L H I Q K D N P K I V H A F D M E
241 atgataaaccttccacatccagaaagacaaccccaagatagtagcatgcttttgacatggag  300
    D L G D K A V Y C R C W R S K K F P F C
301 gatttgggagataaagctgtgtactgccgttggtggagggtccaaaaagttcccattctgt  360
    D G A H T K H N E E T G D N V G P L I I
361 gatgggggctcacacaaaacataacgaagagactggagacaatgtggggccctctgatcatc  420
    K K K E T * SEQ ID NO:625
421 aagaaaaaagaaacttaaattggacacttttgatgctgcaaatacagcttgctcgtgaagtta  480
481 cctgattgtttaattagaatgactaccacctctgtctgattcaccttcgctggattctaa  540
541 atgtggtatattgcaaactgcagctttcacatttatggcatttgtcttggtgaaacatcg  600
601 tgggtgcacatttgttttaacaaaaaaaaaaaaaaaaa SEQ ID NO:624  636
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Figure 2

1 50
MSMT SSVR VEWIAAVTIA AGTAAIGYLA YKRFYVKDHR NKSMINPHIQ
MSLTSSSSVR VEWIAAVTIA AGTAAIGYLA YKRFYVKDHR NKAMINLHIQ
MGLSSNSAVR VEWIAAVTFA AGTAALGYLA YKKFYAKENR TKAMVNLQIQ

51 100
KDNPKVVHAF DMEDLGDKAV YCRCWRSKKF PLCDGSHTKH NEETGDNVGP
KDNPKIVHAF DMEDLGDKAV YCRCWRSKKF PFCDGAHTKH NEETGDNVGP
KDNPKVVHAF DMEDLGDKAV YCRCWRSKKF PFCDGAHIKH NEETGDNVGP

101
LIIKKKDT SEQ ID NO:626 BOVINE
LIIKKKET SEQ ID NO:625 HUMAN
LIIKKKET SEQ ID NO:627 MURINE

Figure 3

1 50
MGLSSNSAVR VEWIAAVTFA AGTAALGYLA YKKFYAKENR TKAMVNLQIQ

51 100
KDNPKVVHAF DMEDLGDKAV YCRCWRSKKF PFCDGAHIKH NEETGDNVGP

101
LIIKKKET SEQ ID NO:627 MURINE